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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/930,283A / DATE: 12/03/2001

TIME: 15:50:27

Input Set : N:\Crf3\RULE60\09930283A.txt
 Output Set: N:\CRF3\12032001\I930283A.raw

SEQUENCE LISTING

4 (1) GENERAL INFORMATION:

- 6 (i) APPLICANT: Kasid, Usha
 Gokhale, Prafulla
 Dritschilo, Anatoly
 Rahman, Aqulur
- 11 (ii) TITLE OF INVENTION: Liposomes containing Oligonucleotides
- 13 (iii) NUMBER OF SEQUENCES: 3
- 15 (iv) CORRESPONDENCE ADDRESS:
 (A) ADDRESSEE: Hendricks and Assoc.
 (B) STREET: P.O. Box 2509
 (C) CITY: Fairfax
 (D) STATE: VA
 (E) COUNTRY: US
 (F) ZIP: 22031

- 23 (v) COMPUTER READABLE FORM:
 (A) MEDIUM TYPE: Floppy disk
 (B) COMPUTER: IBM PC compatible
 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 (D) SOFTWARE: PatentIn Release #1.0, Version #1.25

- 29 (vi) CURRENT APPLICATION DATA:
 (A) APPLICATION NUMBER: US/09/930,283A

- C--> 30 (B) FILING DATE: 16-Aug-2001
- C--> 31 (C) CLASSIFICATION:

- 32 (vii) PRIOR APPLICATION DATA:
 (A) APPLICATION NUMBER: 09/354,109
 (B) FILING DATE: 1999-07-15

- 38 (viii) ATTORNEY/AGENT INFORMATION:
 (A) NAME: Hendricks, Glenna
 (B) REGISTRATION NUMBER: 32,535
 (C) REFERENCE/DOCKET NUMBER: kasid

- 43 (ix) TELECOMMUNICATION INFORMATION:
 (A) TELEPHONE: (703) 591-4470
 (B) TELEFAX: (703) 591-4428

48 (2) INFORMATION FOR SEQ ID NO: 1:

- 50 (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 15 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: unknown
- 56 (ii) MOLECULE TYPE: DNA (genomic)
- 58 (iii) HYPOTHETICAL: NO
- 60 (iv) ANTI-SENSE: YES
- 64 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
 GTGCTCCATT GATGC

68 (2) INFORMATION FOR SEQ ID NO: 2:

- 70 (i) SEQUENCE CHARACTERISTICS:

ENTERED

15

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71 (A) LENGTH: 25 base pairs
72 (B) TYPE: nucleic acid
73 (C) STRANDEDNESS: single
74 (D) TOPOLOGY: unknown
76 (iii) MOLECULE TYPE: DNA (genomic)
78 (iii) HYPOTHETICAL: NO
80 (iv) ANTI-SENSE: YES
84 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
86 CCTGTATGTG CTCCATTGAT GCAGC 25
88 (2) INFORMATION FOR SEQ ID NO: 3:
90 (i) SEQUENCE CHARACTERISTICS:
91 (A) LENGTH: 15 base pairs
92 (B) TYPE: nucleic acid
93 (C) STRANDEDNESS: single
94 (D) TOPOLOGY: unknown
96 (iii) MOLECULE TYPE: DNA (genomic)
98 (iii) HYPOTHETICAL: NO
100 (iv) ANTI-SENSE: YES
104 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
106 GCATCAATGG AGCAC 15

VERIFICATION SUMMARY

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L:30 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]

L:31 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]